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RESULT 3
JC5894
Killer cell inhibitory receptor p91A prec.
C;Species: Mus musculus (house mouse)
C;Date: 18-Mar-1998 #sequence_revision 18
C;Accession: JC5894
R;Yamashita, Y; Fukuta, D; Tsuji, A.; N
J. Biochem. 123, 358-368, 1998
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JC5895
killer cell
C;Species: M
C;Date: 18-M
C;Accession:
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B;Yamashita, Y; Fukuta, D.; Tsuji, A.; Nagabukuro, J. Biochem. 123, 358-368, 1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;1-23/Domain: signal sequence #status predicted <SIG>;24-118,119-220,221-315,316-418,419-517,518-618/Domain:;24-118,10-220, 221-315,316-418,419-517,518-618/Domain: transmembrane #status predicted <TMM>;675-680/Domain: cytoplasmic #status predicted <CYT>
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Mus musculus (house mouse)
                                                                                                                                                              QDHTMENLIRMGMAVVVFIVLSILATEAWRSHRQTHCAAGN
                                                                                                                                                                                     QDYTVENLIRMGVAGLVLVVLGILLFEAQHSQRSLQDAAGS
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                                                                                                                                                                                                                                                                                 LPLTPSLSVQPNHTVHSGETVSLLCWSMDSVDTFILSKEGSAQQPLRLKSKSHDQQSQAE
                                                                                                                                                                                                                                                      FRMGPVTSAHVGTYRCYSSLSSNPYLLSLPSDPLELVVSASLGQ----
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Pred. No. 1.4e-68;
                                                                                      precursor
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                          Nagabukuro,
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                         A.; Matsuda,
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JC5896
killer cell inhibitory receptor p91C precursor - mouse
C:Species: Mus musculus (house mouse)
C;Date: 18-Mar-1998 #sequence_revision 18-Mar-1998 #text_change (
C;Accession: JC5896
R;Yamashita, Y:; Fukuta, D:; Tsuji, A.; Nagabukuro, A.; Matsuda,
J. Biochem. 123, 358-368, 1998
J. Biochem. 123, 358-368, 1998
A;Residues: 1-635 <YAM>
A;Cross-references: GB:/
C;Comment: This protein
                                          A;Accession: JC5896
A;Status: nucleic aci
A;Molecule type: mRNA
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A; Reference number: JC5894;
                                                        ;Status: nucleic acid sequence
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A; Residues: 1-841 <Y;
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A;Reference number: JC5894;
A;Accession: JC5894
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                         ORGOCILORGGASGT
                                                GLH----MYLKALIGVSVAFILFLFILIFILLRRRHRGKFRKDVQKEKDLQLSSGAEEPIT
                                                                          GQHPQDYTVENLIRMGVAG-LVLVVLGILLFEAQH-----SQRSLQDAAGSEQ---
                                                                                                       SQAEFSMSAVTSHLSGTYRCYGAQNSSFYLLSSASAPVELTVSGPIETSTPPPTMSMPLG
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45.9%; Pre
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Pred. No. 1.9e-64;
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n: extracellular Ig-like
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GB:AF041036; tein function

NID:92791693; PIDN:AAB96928.1; PID:92791694 as ingibitory cell-surface molecule against

against

cell

ructures and chromosoma JC5894; MUID:98218758;

chromosomal

PMID:9538215

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novel

murine Nishikawa,

regulatory

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05-Nov-1999

not

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W09848017-A1

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This sequence represents a novel leukocyte immunoglobulin-like receptor (LIR) polypeptide LIR-pmb25. This sequence can be administered therapeutically to treat disorders associated with insufficient/defective amounts of LIR polypeptide. LIR-p3G2 and certain other LIR family members contain cytoplasmic immunoreceptor tyrosine-based inhibitory members (ITIMs), whilst other LIR family members lack ITIMs. By analogy with the structure and function of known MHC Class I receptor molecules, LIRs whilst those lacking ITIMs are activatory receptors mediating negative signalling, receptor that mediates negative signalling could result in autoimmune could result in suppressed immune function. They are also useful to char encodes for detecting LIR mucleic acids or isolating LIR DNA from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Leukocyte immunoglobulin-like receptor, LIR, polypeptides - useful, for treating autoimmune diseases or disease states associated with suppressed immune function.
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                              AS
                                                                  GAGHPPLHLRSEHQAQQNQAEFRMGPVTSAHVGTYRCYSSLSSNPYLLSLPSDPLELVVS
                                                                                                                                HNLSSEWSAPSDPLDILITGOFYDRPSLSVOPVPTVAPGKNVTLLCOSRGOFHTELLTKE
                                                   GAADSPLRLKSKROSHKYQAEFPMSPVTSAHAGTYRCYGSLSSNPYLLTHPSDPLELVVS
                                                                                                               YNLSSEWSAPSDPLDILITGQIRARPFLSVRPGPTVASGENVTLLCQSQGGMHTFLLTKE
                                                                                                                                                                               KLTFQCGSDAGYDRFVLYKEWGRDFLQRPGRQPQAGLSQANFTLGPVSRSYGGQYTCSGA
                                                                                                                                                                                                   SLTLQCVSDVGYDRFVLYKEGERDFLQRPGWQPQAGLSQANFTLGPVSPSHGGQYRCYSA
                                                                                                                                                                                                                                                                    IPSVGPVSPSRRWSYRCYAYDSNSPYVWSLPSDLLELLVPGVSKKPSLSVQPGPMVAPGE
                                                                                                                                                                                                                                                                                                         AYSKPTLSALPSPVVTSGRNVTIQCDSQVAFDGFILCKEGEDEHPQCLNSHSHARGSSRA
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                                                                                                                                                                                                                                             IFSVGÞVSÞSRRWSYRCYGYDSRAÞYVWSLÞSDLLGLLVÞGVSKXÞSLSVÓÞGÞVVAÞGE
                                                                                                                                                                                                                                                                                                                               AYSKPTLSALPSPVVTLGGNVTLQCVSQVAFDGFILCKEGEDEHPQRLNSHSHARGWSWA
                                                                                                                                                                                                                                                                                                                                                              YRENKSASWVRRI-QEPGKNGQFPIPSITWEHAGRYHCQYYSHNHS-SEYSDPLELVVTG
                                                                                                                                                                                                                                                                                                                                                                                                                                      MTPILTVLICLGLSLDPRTHVQAGPLPKPTLWAEPGSVITQGSPVTLRCQGSLETQEYHL
                              420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63.7%;
81.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1754; DB 2;
Pred. No. 5.4e-126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 439;
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Best Local Simi
Matches 343;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents human gp49 polypeptide HM43. The present invention also describes: (1) a fusion polypeptide comprising a human gp49 component and FceRI or a detectable marker; (2) a recombinant nucleic acid encoding human gp49 or a human gp49-FceRI fusion as above; (3) a cell or vector comprising the recombinant nucleic acid as in (2), and (4) an antibody which selectively binds to gp49. Mammalian gp49 or its related DNA can be used to treat an undesired immune response, especially a mast cell-related disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 439 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 5; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cell-surface member of immunoglobulin super-family, human gp49 - 1 to treat undesired immune responses, especially mast cell-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-193318/17
N-PSDB; AAV23274.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-MAR-1998.
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                                                                                                                                                                                                                                      121
                                                                                                                                                                                                                                                                                          119
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arrow; cell-surface member; FceRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BRIGHAM &
KLTFQCGSDAGYDRFVLYKEWGRDFLQRPGRQPQAGLSQANFTLGPVSRSYGGQYTCSGA
                                  SLTLQCVSDVGYDRFVLYKEGERDFLQRPGWQPQAGLSQANFTLGPVSPSHGGQYRCYSA 298
                                                                                                                                             IFSVGPVSPSRRWSYRCYAYDSNSPYVWSLPSDLLELLVPGVSKKPSLSVQPGPMVAPGE
                                                                                                                                                                                                                        AYSKPTLSALPSPVVTSGGNVTIQCDSQVAFDGFILCKEGEDEHPQCLNSHSHARGSSRA
                                                                                                                                                                                                                                                             AYSKPTLSALPSPVVTLGGNVTLQCVSQVAFDGFILCKEGEDEHPQRLNSHSHARGWSWA
                                                                                                                                                                                                                                                                                                                                                                        YRENKSASWVRRI-QEPGKNGQFPIPSITWEHAGRYHCQYYSHNHS-SEYSDPLELVVTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MTPILTVLICLGLSLGPRTHVQAGHLPKPTLWAEPGSVIIQGSPVTLRCQGSLQAEEYHL
                                                                                                                                                                                                                                                                                                                                  YREKKTALWITRIPQELVKKGQFPILSITWEHAGRYCCIYGSHTAGLSESSDPLELVVTG
                                                                                                                                                                                                                                                                                                                                                                                                                                             MTSILTVLICLGLSLDPRTHVQAGPLPRPTLWAEPGSVITQGSPVTLRCQGSLETQBYHL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       63.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 1754; DI
Pred. No. 5.4e-
18; Mismatches
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